



TITLE: METHOD AND APPARATUS FOR
AUTOMATICALLY REMOVING VECTOR UNIT IN
DNA BASE SEQUENCE
INVENTORS: Kensaku IMAI, et al.
SERIAL NO.: 09/785,269
DOCKET NO.: 826.1335C

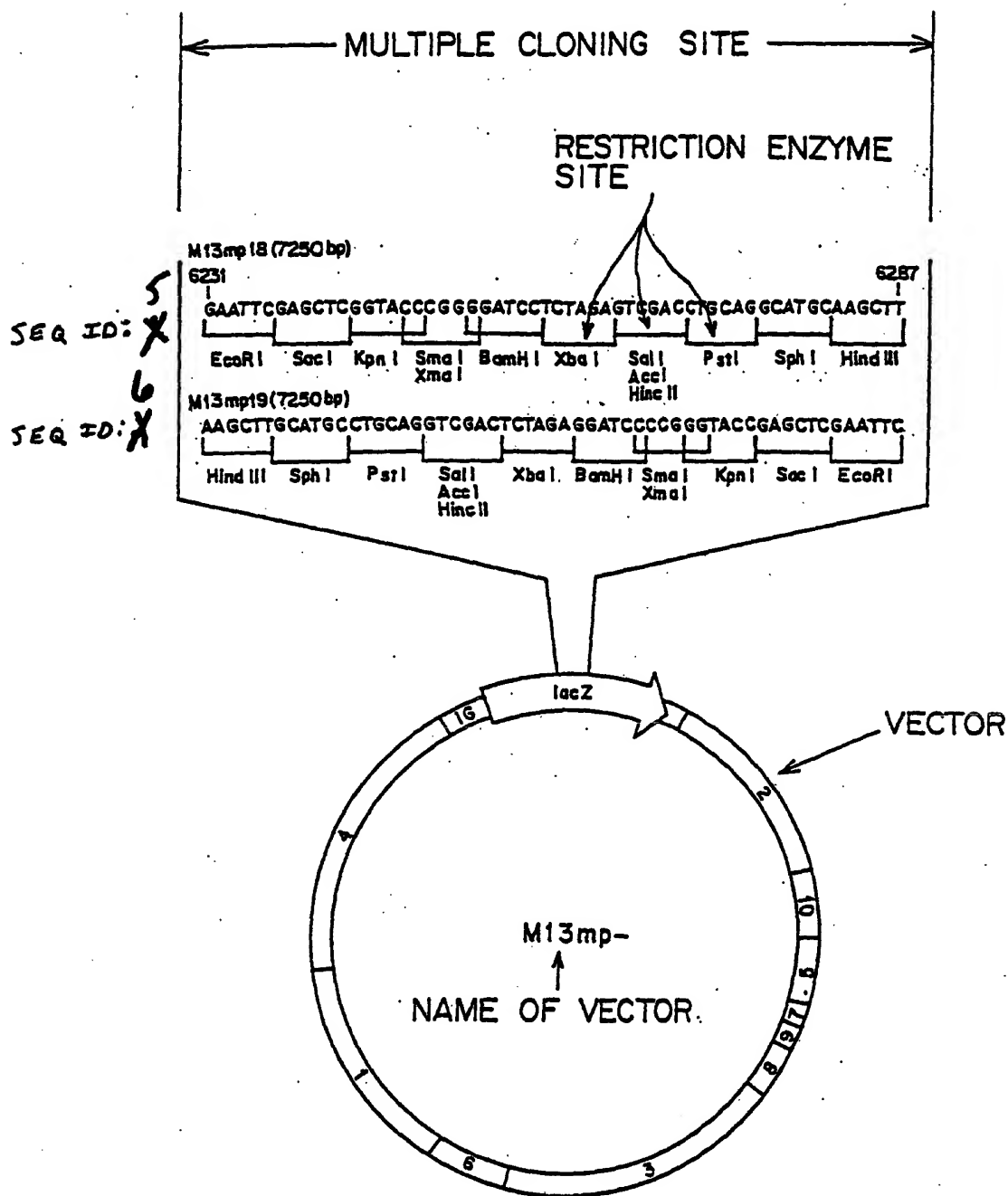


FIG. 3

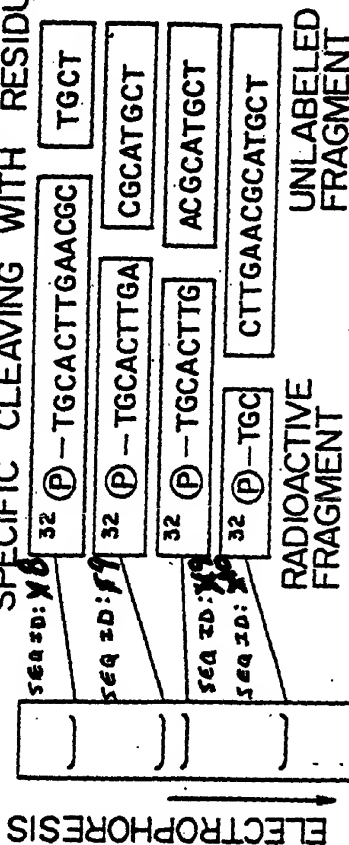


TITLE METHOD AND APPARATUS FOR
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DNA FRAGMENT LABELED WITH ³²P AT 5' EDGE

SEQ ID: 1 ³²P - TGCACCTTGAACGCATGCT

RADIOACTIVE FRAGMENTS OF VARIOUS
LENGTHS THROUGH CHEMICAL PROCESS OF
SPECIFIC CLEAVING WITH RESIDUAL BASE A



THESE FRAGMENTS CAN BE STRICTLY
ISOLATED DEPENDING ON LENGTH
THROUGH GEL ELECTROPHORESIS

FIG. 4



VECTOR DB FORMAT

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>ID
PUC18
>SEQ ID: X //
TCGCGCGTTTCGGTGATGACGGTGAAAACTCTGACACATGACGCTCCCGGAGACGGTCACAGCTTGTCTGTAAGCGGAT
GCCGGGAGCAGACAAGCCCGTCAGGGCGCGTCAGCGGGTGTGGCGGGTGTGGGGCTGGCTTAACATATGCGGCATCAGA
GCAGATTGTACTGAGAGTGACCATATGCGGTGTGAAATACCGCACAGATGCGTAAGGAGAAAATACCGCATCAGGCGCC
ATTCGCCATTGAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCTCTTCGCTATTACGCCAGCTGGCGAAAGGG
GGATGTGCTGCAAGGCGATTAAAGTTGGGTAAACGCCAGGGTTTTCCAGTCACGACGTTGTAAAACGACGGCCAGTGCCAA
GCTTGCACTGCCTGCGAGGTCGACTCTAGAGGATCCCCGGGTACCGAGCTCGAATTCGTAATCATGGTCATAGCTGTTTCT
GTGTGAAATTGTTATCCGCTCACAATCCACACAACATACGAGCCGGAAGCATAAAGTGTAAAGCCTGGGGTGCCTAATG
AGTGAGCTAACTCACATTAAATGCGTTGCGCTCACTGCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCATTAA
GAATCGGCCAACGCGCGGGGAGAGGCGGTTTTCGCTATTGGGCGCTCTTCGCTTCTCGCTCACTGACTCGCTGCGCTCG
GTCGTTCCGCTGCGGCGAGCGGTATCAGCTCAAGAGCGGTAATACGGTTATCCACAGAATCAGGGGATAACGCAGG
AAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCTTGTGGCGTTTTTCCATAGGCTCC
GCCCCCTGACGAGCATCAGAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAGATACAGGGCG
TTTTCCCTGGAAAGCTCCCTCGTGGCTCTCTGTTCCGACCCCTGCCGCTTACCGGATACCTGTCCGCTTTCTCCCTTC
GGGAAGCGTGGCGCTTTCTCAAGCTCAGCTGTAGGTATCTCAGTTGCGGTGTAGGTGCTTCGCTCCAAAGCTGGGCTGTG
TGCAGGAACCCCGCTTCCGCGGACCGCTGCGCTTATCCGTTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGAC
TTATCGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTTGAAGTG
GTGGCCTAACTACGGCTACACTAGAAGAACAGTATTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGAAAAAGAG
TTGGTAGCTCTTGATCCGGCAAAACAAACCCGCTGGTAGCGGTGGTTTTTTGTTTGAAGCAGCAGATTACCGCGAGA
AAAAAGGATCTCAAGAAGATCCTTTGATCTTTCTACGGGCTCTGACGCTCAGTGAACGAAAACTCACGTTAAGGGAT
TTTGGTCATGAGATTATCAAAAAGGATCTTCACCTAGATCCTTTAAATTAAGTGAAGTTTTAAATCAATCTAAAGTA
TATATGAGTAACTTGGTCTGACAGTTACCAATGCTTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTCGTTCA
TCCATAGTTGCTGACTCCCGCTCGTGTAGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGAT
ACCGCGAGACCCAGCTCACCAGCTCCAGATTTATCAGCAATAAACCAGCCAGCCGGAAGGGCCGAGCGCAGAGTGGTC
CTGCAACTTTATCCGCTCCATCCAGTCTATTAAATGTTGCGGGAAAGCTAGAGTAAGTAGTTCCGCCAGTTAATAGTTTG
CGCAACGTTGTTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCTGTTGGTATGGCTTCATTAGCTCCGTTCCCA
ACGATCAAGGCGAGTTACATGATCCCCATGTTGTGCAAAAAGCGGTTAGCTCCTTCGCTCCGATCGTTGTCAGAA
GTAAGTTGGCCGAGTGTATCACTCATGGTTATGGCAGCACTGCATAATTCTTACTGTATGCCATCCGTAAGATGC
TTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTGAGAATAGTGTATCGGCGACCGAGTTGCTCTTGGCCGCGTC
AATACGGGATAATACCGCGCCACATAGCAGAAGTTAAAAGTGCTCATATTGGAAAACTGTTCTCGGGGCGAAAACTCT
CAAGGATCTTACCGCTGTTGAGATCCAGTTCGATGTAACCCACTCGTGACCCAACTGATCTTCAGCATCTTTACTTTC
ACCAGCGTTTCTGGGTGAGCAAAAACAGGAAGGCAAAATGCCGCAAAAAGGGAAATAAGGGCGACACGGAAATGTTGAAT
ACTCATACTCTTCTTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATATTTGAATGTA
TTAGAAAAATAAACAAATAGGGGTTCCGCGCACATTTCCCGAAAAAGTGCCACCTGACGTCTAAGAAACCATTTATC
ATGACATTAACCTATAAAAAATAGGCGTATCACGAGGCCCTTTCGTC
>MULTI
399.. 450
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FIG. 9



THE METHOD AND APPARATUS FOR
AUTOMATICALLY REMOVING VECTOR UNIT IN
DNA BASE SEQUENCE

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(* INDICATES MULTIPLE CLONING SITE)

¹² SEQ ID: ~~1~~ GTGCCAAGCTTGCATGCCCTGCAGTCGACTCTAGAGGATCCCCGGGTACCGAGCTCGAATTCGTAAT

SEQ ID: ¹³ ~~14~~ AAGCTT⇒HIND III

SEQ ID: ~~14~~ ¹⁵ GCATGC⇒SPH I

SEQ ID: ~~15~~ ¹⁶ CTGCAG⇒PST I

SEQ ID: ~~16~~ ¹⁷ GTCGAC⇒SAL I, ACC I, HINC II

SEQ ID: ~~17~~ ¹⁸ TCTAGA⇒XBA I

SEQ ID: ~~18~~ ¹⁹ GGATCC⇒BAMH I

SEQ ID: ~~19~~ ²⁰ CCCGGG⇒SMA I, XMA I

SEQ ID: ~~20~~ ²¹ GGTACC⇒KPN I

SEQ ID: ~~21~~ ²² GAGCTC⇒SAC I

SEQ ID: ~~22~~ ²³ GAATTC⇒ECOR I

FIG. 10

01
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WHEN HIND III IS SPECIFIED ON VECTOR 5' SIDE
XBA I IS SPECIFIED ON VECTOR 3' SIDE, HIND III IS
SPECIFIED ON OBJECT DNA 5' SIDE, AND XBA I IS
SPECIFIED ON OBJECT DNA 3' SIDE

(**** INDICATES RESIDUAL MULTIPLECLONING SITE
(++++ INDICATES AN OBJECT DNA FRAGMENT

(SEQUENCE
ID NO. 4)

GTGCCAAGCTT+++++
AAGCTT
↑

5' SIDE RETRIEVAL KEY
(IN THIS EXAMPLE,
HIND III SITE)

(SEQUENCE
ID NO. 23)

TCTAGAGGATCCCCGGTACCGAGCTCGAATTCGTAAT
TCTAGA
↑

9' SIDE RETRIEVAL KEY
(IN THIS EXAMPLE, XBA I SITE)

FIG. 17